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MORI, Yukiko
ITO, Hisao

<120> CARBAMOYL-PHOSPHATE SYNTHETASE GENE OF CORYNEFORM BACTERIA AND METHOD FOR PRODUCING L-ARGININE

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Glu	Ala	Lys	Phe 500	Met	Gly	Leu	Ser	Asp 505	Leu	Gln	Ile	Ala	Ala 510	Leu	Arg
Pro	Glu	Phe 515	Ala	Gly	Glu	Asp	Gly 520	Val	Arg	Thr	Leu	Arg 525	Leu	Ser	Leu
Gly	Ile 530	Arg	Pro	Val	Phe	Lys 535	Thr	Val	Asp	Thr	Cys 540	Ala	Ala	Glu	Phe
Glu 545	Ala	Lys	Thr	Pro	Tyr 550	His	Tyr	Ser	Ala	Tyr 555	Glu	Leu	Asp	Pro	Ala 560
Ala	Glu	Ser	Glu	Val 565	Ala	Pro	Gln	Thr	Glu 570	Arg	Glu	Lys	Val	Leu 575	Ile
Leu	Gly	Ser	Gly 580	Pro	Asn	Arg	Ile	Gly 585	Gln	Gly	Ile	Glu	Phe 590	Asp	Tyr
Ser	Cys	Val 595	His	Ala	Ala	Leu	Glu 600	Leu	Ser	Arg	Val	Gly 605	Tyr	Glu	Thr
Val	Met 610	Val	Asn	Cys	Asn	Pro 615	Glu	Thr	Val	Ser	Thr 620	Asp	Tyr	Asp	Thr
Ala 625	Asp	Arg	Leu	Tyr	Phe 630	Glu	Pro	Leu	Thr	Phe 635	Glu	Asp	Val	Met	Glu 640
Val	Tyr	His	Ala	Glu 645	Ala	Gln	Ser	Gly	Thr 650	Val	Ala	Gly	Val	Ile 655	Val
Gln	Leu	Gly	Gly 660	Gln	Thr	Pro	Leu	Gly 665	Leu	Ala	Asp	Arg	Leu 670	Lys	Lys
Ala	Gly	Val 675	Pro	Val	Ile	Gly	Thr 680	Ser	Pro	Glu	Ala	Ile 685	Asp	Met	Ala
Glu	Asp 690	Arg	Gly	Glu	Phe	Gly 695	Ala	Leu	Leu	Asn	Arg 700	Glu	Gln	Leu	Pro
Ala 705	Pro	Ala	Phe	Gly	Thr 710	Ala	Thr	Ser	Phe	Glu 715	Glu	Ala	Arg	Thr	Val 720

4st

Ala Asp Glu Ile Ser Tyr Pro Val Leu Val Arg Pro Ser Tyr Val Leu 725 730 735

Gly Gly Arg Gly Met Glu Ile Val Tyr Asp Glu Ala Ser Leu Glu Asp 740 745 750

Tyr Ile Asn Arg Ala Thr Glu Leu Ser Ser Asp His Pro Val Leu Val 755 760 765

Asp Arg Phe Leu Asp Asn Ala Ile Glu Ile Asp Val Asp Ala Leu Cys 770 780

Asp Gly Asp Glu Val Tyr Leu Ala Gly Val Met Glu His Ile Glu Glu 785 790 795 800

Ala Gly Ile His Ser Gly Asp Ser Ala Cys Ala Leu Pro Pro Met Thr 805 810 815

Leu Gly Ala Gln Asp Ile Glu Lys Val Arg Glu Ala Thr Lys Lys Leu 820 825 830

Ala Leu Gly Ile Gly Val Gln Gly Leu Met Asn Val Gln Tyr Ala Leu 835 840 845

Lys Asp Asp Ile Leu Tyr Val Ile Glu Ala Asn Pro Arg Ala Ser Arg 850 855 860

Thr Val Pro Phe Val Ser Lys Ala Thr Gly Val Asn Leu Ala Lys Ala 865 870 875 880

Ala Ser Arg Ile Ala Val Gly Ala Thr Ile Lys Asp Leu Gln Asp Glu 885 890 895

Gly Met Ile Pro Thr Glu Tyr Asp Gly Gly Ser Leu Pro Leu Asp Ala 900 905 910

Pro Ile Ala Val Lys Glu Ala Val Leu Pro Phe Asn Arg Phe Arg Arg 915 920 925

Pro Asp Gly Lys Thr Leu Asp Thr Leu Leu Ser Pro Glu Met Lys Ser 930 935 940

Thr Gly Glu Val Met Gly Leu Ala Asn Asn Phe Gly Ala Ala Tyr Ala 945 950 955 960

Lys Ala Glu Ala Gly Ala Phe Gly Ala Leu Pro Thr Glu Gly Thr Val 965 970 975

Phe Val Thr Val Ala Asn Arg Asp Lys Arg Thr Leu Ile Leu Pro Ile 980 985 990

Gln Arg Leu Ala Ser Met Gly Tyr Lys Ile Leu Ala Thr Glu Gly Thr 995 1000 1005

Ala Gly Met Leu Arg Arg Asn Gly Ile Asp Cys Glu Val Val Leu 1010 1015 1020

Lys Ala Ser Asp Ile Arg Glu Gly Val Glu Gly Lys Ser Ile Val 1025 1030 1035

Asp Arg Ile Arg Glu Gly Glu Val Asp Leu Ile Leu Asn Thr Pro 1040 1045 1050

Ala Gly Ser Ala Gly Ala Arg His Asp Gly Tyr Asp Ile Arg Ala 1055 1060 1065

Ala Ala Val Thr Val Gly Val Pro Leu Ile Thr Thr Val Gln Gly 1070 1075 1080

Val Thr Ala Ala Val Gln Gly Ile Glu Ala Leu Arg Glu Gly Val 1085 1090 1095

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